



ERP01.003APC.TXT

SEQUENCE LISTING

<110> Schor, Seth Lawrence
Schor, Ana Maria

<120> Polypeptides, Polynucleotides and Uses
Thereof

<130> ERP01.003APC

<140> 09/581,651
<141> 2000-10-10

<150> PCT/GB98/03766
<151> 1998-12-15

<150> GB 9726539.1
<151> 1997-12-16

<160> 45

<170> FastSEQ for Windows Version 4.0

<210> 1
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<212> PRT
<213> Homo sapiens

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Leu Asn Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Leu Ala Val
20 25 30
Leu Cys Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys
35 40 45
Arg Gln Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser
50 55 60
Gln Ser Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn
65 70 75 80
Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys
85 90 95
Tyr Gly Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu
100 105 110
Glu Thr Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp
115 120 125
Thr Tyr Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile
130 135 140
Gly Ala Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His
145 150 155 160
Glu Gly Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His
165 170 175
Glu Thr Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys
180 185 190
Gly Glu Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala
195 200 205
Ala Gly Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln
210 215 220
Gly Trp Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg
225 230 235 240
Ile Thr Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr
245 250 255
Ser Tyr Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn

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Leu	Leu	Gln	Cys	Ile	Cys	Thr	Gly	Asn	Gly	Arg	Gly	Glu	Trp	Lys	Cys
260						265						270			
275						275	280					285			
Glu	Arg	His	Thr	Ser	Val	Gln	Thr	Thr	Ser	Ser	Gly	Ser	Gly	Pro	Phe
290						290	295					300			
Thr	Asp	Val	Arg	Ala	Ala	Val	Tyr	Gln	Pro	Gln	Pro	His	Pro	Gln	Pro
305						310						315			320
Pro	Pro	Tyr	Gly	His	Cys	Val	Thr	Asp	Ser	Gly	Val	Val	Tyr	Ser	Val
325															
Gly	Met	Gln	Trp	Leu	Lys	Thr	Gln	Gly	Asn	Lys	Gln	Met	Leu	Cys	Thr
340															
Cys	Leu	Gly	Asn	Gly	Val	Ser	Cys	Gln	Glu	Thr	Ala	Val	Thr	Gln	Thr
355															
Tyr	Gly	Gly	Asn	Ser	Asn	Gly	Glu	Pro	Cys	Val	Leu	Pro	Phe	Thr	Tyr
370															
Asn	Gly	Arg	Thr	Phe	Tyr	Ser	Cys	Thr	Thr	Glu	Gly	Arg	Gln	Asp	Gly
385															400
His	Leu	Trp	Cys	Ser	Thr	Thr	Ser	Asn	Tyr	Glu	Gln	Asp	Gln	Lys	Tyr
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Ser	Phe	Cys	Thr	Asp	His	Thr	Val	Leu	Val	Gln	Thr	Gln	Gly	Gly	Asn
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Ser	Asn	Gly	Ala	Leu	Cys	His	Phe	Pro	Phe	Leu	Tyr	Asn	Asn	His	Asn
435															
Tyr	Thr	Asp	Cys	Thr	Ser	Glu	Gly	Arg	Arg	Asp	Asn	Met	Lys	Trp	Cys
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Gly	Thr	Thr	Gln	Asn	Tyr	Asp	Ala	Asp	Gln	Lys	Phe	Gly	Phe	Cys	Pro
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Met	Ala	Ala	His	Glu	Glu	Ile	Cys	Thr	Thr	Asn	Glu	Gly	Val	Met	Tyr
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Arg	Ile	Gly	Asp	Gln	Trp	Asp	Lys	Gln	His	Asp	Met	Gly	His	Met	Met
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545															
Cys	Thr	Cys	Phe	Gly	Gln	Gly	Arg	Gly	Arg	Trp	Lys	Cys	Asp	Pro	Val
565															
Asp	Gln	Cys	Gln	Asp	Ser	Glu	Thr	Gly	Thr	Phe	Tyr	Gln	Ile	Gly	Asp
580															
Ser	Trp	Glu	Lys	Tyr	Val	His	Gly	Val	Arg	Tyr	Gln	Cys	Tyr	Cys	Tyr
595															
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625															
Pro	Asn	Ser	His	Pro	Ile	Gln	Trp	Asn	Ala	Pro	Gln	Pro	Ser	His	Ile
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<213> Homo sapiens

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atcaacagtg	ggagcggacc	tacctaggca	atgcgttggt	ttgtacttgt	tatggaggaa	300
gccgaggtt	taactgcgag	agtaaacctg	aagctgaaga	gacttgctt	gacaagtaca	360
ctggAACAC	ttaccgagtg	ggtgcacatt	atgagcgtcc	taaagactcc	atgatctggg	420
actgtaccc	catcggggct	gggcgaggga	gaataagctg	taccatcgca	aaccgctgcc	480
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gttacatgtt	agagtgtgt	tgtcttggta	atggaaaagg	agaatggacc	tgcaagccca	600
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agaagcccta	ccaaggctgg	atgatggtag	attgtacttg	cctggggagaa	ggcagcggac	720
gcatcactg	cacttctaga	aatagatgca	acgatcagga	cacaaggaca	tcctataagaa	780
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gatctggccc	cttcaccgt	gttcgtcag	ctgtttacca	accgcagct	caccccccagc	960
ctcctcccta	tggccactgt	gtcacagaca	gtgggtgtgt	ctactctgtg	gggatgcagt	1020
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gccaagagac	agctgttaacc	cagacttacg	gtggcaactc	aatggagag	ccatgtgtct	1140
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aatactctt	ctgcacagac	cacactgttt	tggttcagac	tcgaggagga	aattccaatg	1260
gtgccttgc	ccacttcccc	ttcctataca	acaaccacaa	ttacactgtat	tgcacttctg	1320
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agaagtatgt	gcatgggtgc	agataccatg	gctactgcta	tggccgtggc	attggggagt	1800
ggcattgcca	acctttacag	acctatccaa	gctcaagtgg	tcctgtcgaa	gtatttatca	1860
ctgagactcc	gagtcaagccc	aactcccacc	ccatccagtg	gaatgcacca	cagccatctc	1920
acatttccaa	gtacattctc	aggtggagac	ctgtgagtat	cccacccaga	aacttggat	1980
actgagtctc	ctaattttat	caattctgtat	ggtttctttt	tttcccagct	tttgagccaa	2040
caactctgtat	taactattcc	tatagcattt	actatatttg	tttagtgaac	aaacaatatg	2100
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<400> 3
 Ile Ser Lys Tyr Ile Leu Arg Trp Arg Pro Val Ser Ile Pro Pro Arg
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 Asn Leu Gly Tyr
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<210> 4
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 Tyr Gly Gly Ser Arg
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Ser Asn Tyr Glu Gln Asp Gln
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Ala Leu Cys His
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1 5 10 15
Arg Asp Gln Cys Ile
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<211> 21
<212> PRT
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1 5 10 15
Tyr Gly Gly Ser Arg
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Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Gly Arg Thr Phe Tyr Ser
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Ser Asn Tyr Glu Gln Asp Gln
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<213> Homo sapiens

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Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn Ser Asn
1 5 10 15
Gly Ala Leu Cys His
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<210> 11
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<213> Homo sapiens

<400> 11
Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Thr Ala Tyr Ser Gln Leu
1 5 10 15
Arg Asp Gln Cys Ile
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<210> 12
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<213> Homo sapiens

<400> 12
Ile Ser Lys Thr Ile Leu Arg Trp Arg Pro Lys Asn Ser Val Gly Arg
1 5 10 15
Trp Lys Glu Ala
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<210> 13
<211> 18
<212> PRT
<213> Homo sapiens

<400> 13
Asn Leu Val Ala Thr Cys Leu Pro Val Arg Ala Ser Leu Pro His Arg
1 5 10 15
Leu Asn

<210> 14
<211> 31
<212> PRT
<213> Homo sapiens

<400> 14
Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Ala Val Gln Cys
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Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys Arg
20 25 30

<210> 15
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<213> Homo sapiens

<400> 15
Gln Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser Gln
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Ser Lys Pro Gly
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<212> PRT
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<400> 16
Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn Gln Gln Trp Glu Arg
1 5 10 15
Thr Tyr Leu Gly Asn Ala Leu Val Cys Thr Cys Tyr Gly Gly Ser Arg
20 25 30
Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu Glu Thr
35 40 45

<210> 17
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<400> 17
Cys Asn Asp Gln Asp Thr Arg Thr Ser Tyr Arg Ile Gly Asp Thr Trp
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Ser Lys Lys Asp Asn Arg Gly Asn Leu Leu Gln Cys Ile Cys Thr Gly
20 25 30
Asn Gly Arg Gly Glu Trp Lys Cys Glu Arg
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<210> 18
<211> 35
<212> PRT
<213> Homo sapiens

<400> 18
His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe Thr Asp
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Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro Pro Pro
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Tyr Gly His
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<210> 19
<211> 37
<212> PRT
<213> Homo sapiens

<400> 19
Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val Gly Met Gln Trp Leu
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20 25 30
Val Ser Cys Gln Glu
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<210> 20
<211> 45
<212> PRT

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<213> Homo sapiens

<400> 20

Thr Ala Val Thr Gln Thr Tyr Gly Gly Asn Ser Asn Gly Glu Pro Cys
1 5 10 15
Val Leu Pro Phe Thr Tyr Asn Asp Arg Thr Asp Ser Thr Thr Ser Asn
20 25 30
Tyr Glu Gln Asp Gln Lys Tyr Ser Phe Cys Thr Asp His
35 40 45

<210> 21

<211> 48

<212> PRT

<213> Homo sapiens

<400> 21

Cys Thr Thr Asn Glu Gly Val Met Tyr Arg Ile Gly Asp Gln Trp Asp
1 5 10 15
Lys Gln His Asp Met Gly His Met Met Arg Cys Thr Cys Val Gly Asn
20 25 30
Gly Arg Gly Glu Trp Thr Cys Ile Ala Tyr Ser Gln Leu Arg Asp Gln
35 40 45

<210> 22

<211> 43

<212> PRT

<213> Homo sapiens

<400> 22

Cys Ile Val Asp Asp Ile Thr Tyr Asn Val Asn Asp Thr Phe His Lys
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Arg His Glu Glu Gly His Met Leu Asn Cys Thr Cys Phe Gly Gln Gly
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Arg Gly Arg Trp Lys Cys Asp Pro Val Asp Gln
35 40

<210> 23

<211> 48

<212> PRT

<213> Homo sapiens

<400> 23

Cys Gln Asp Ser Glu Thr Gly Thr Phe Tyr Gln Ile Gly Asp Ser Trp
1 5 10 15
Glu Lys Tyr Val His Gly Val Arg Tyr Gln Cys Tyr Cys Tyr Gly Arg
20 25 30
Gly Ile Gly Glu Trp His Cys Gln Pro Leu Gln Thr Tyr Pro Ser Ser
35 40 45

<210> 24

<211> 39

<212> PRT

<213> Homo sapiens

<400> 24

Ser Gly Pro Val Glu Val Phe Ile Thr Glu Thr Pro Ser Gln Pro Asn
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Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His Ile Ser Lys
20 25 30

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Tyr Ile Leu Arg Trp Arg Pro
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<210> 25
<211> 10
<212> PRT
<213> Homo sapiens

<400> 25
Val Ser Ile Pro Pro Arg Asn Leu Gly Tyr
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<210> 26
<211> 44
<212> PRT
<213> Homo sapiens

<400> 26
Trp Phe Leu Phe Phe Pro Ala Phe Glu Pro Thr Thr Leu Ile Asn Tyr
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Ser Tyr Ser Ile Tyr Tyr Ile Cys Leu Val Asn Lys Gln Tyr Val Val
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Asn Ile Asp Leu Thr Glu Lys Lys Lys Lys Lys Lys
35 40

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<211> 6
<212> PRT
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Met Leu Arg Gly Pro Gly
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<210> 28
<211> 65
<212> PRT
<213> Homo sapiens

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Thr Val Leu Val Gln Thr Arg Gly Gly Asn Ser Asn Gly Ala Leu Cys
1 5 10 15
His Phe Pro Phe Leu Tyr Asn Asn His Asn Tyr Thr Asp Cys Thr Ser
20 25 30
Glu Gly Arg Arg Asp Asn Met Lys Trp Cys Gly Thr Thr Gln Asn Tyr
35 40 45
Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro Met Ala Ala His Glu Glu
50 55 60
Ile
65

<210> 29
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<213> Homo sapiens

<400> 29
Val Ser

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<213> Homo sapiens

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Ser Tyr Gln Phe
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Trp Phe Leu Phe Phe Pro Ala Phe Glu Pro Thr Thr Leu Ile Asn Tyr
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Ser Tyr Ser Ile Tyr Tyr Ile Cys Leu Val Asn Lys Gln Tyr Val Val
20 25 30
Asn

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<213> Homo sapiens

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Ile Asp Leu
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<400> 33
Thr Glu Lys Lys Lys Lys Lys Lys
1 5

<210> 34
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<212> PRT
<213> Homo sapiens

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Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Asp Arg Thr Asp Ser Thr
1 5 10 15
Thr Ser Asn Tyr Glu Gln Asp Gln
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<213> Homo sapiens

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Thr Asp His Thr Val Leu Val Gln Thr Arg Gly Gly Asn Ser Asn Gly
1 5 10 15
Ala Leu Cys His
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<210> 36
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<213> Homo sapiens

<400> 36
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Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys Arg Gln
20 25 30
Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser Gln Ser
35 40 45
Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn Gln Gln
50 55 60
Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys Tyr Gly
65 70 75 80
Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu Glu Thr
85 90 95
Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp Thr Tyr
100 105 110
Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile Gly Ala
115 120 125
Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His Glu Gly
130 135 140
Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His Glu Thr
145 150 155 160
Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys Gly Glu
165 170 175
Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala Ala Gly
180 185 190
Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln Gly Trp
195 200 205
Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg Ile Thr
210 215 220
Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr Ser Tyr
225 230 235 240
Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn Leu Leu
245 250 255
Gln Cys Ile Cys Thr Gly Asn Gly Arg Gly Glu Trp Lys Cys Glu Arg
260 265 270
His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe Thr Asp
275 280 285
Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro Pro Pro
290 295 300
Tyr Gly His Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val Gly Met
305 310 315 320
Gln Trp Leu Lys Thr Gln Gly Asn Lys Gln Met Leu Cys Thr Cys Leu
325 330 335
Gly Asn Gly Val Ser Cys Gln Glu Thr Ala Val Thr Gln Thr Tyr Gly
340 345 350
Gly Asn Ser Asn Gly Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Gly
355 360 365
Arg Thr Phe Tyr Ser Cys Thr Thr Glu Gly Arg Gln Asp Gly His Leu
370 375 380
Trp Cys Ser Thr Thr Ser Asn Tyr Glu Gln Asp Gln Lys Tyr Ser Phe

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385 Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn Ser Asn
 390 405 410 415
 Gly Ala Leu Cys His Phe Pro Phe Leu Tyr Asn Asn His Asn Tyr Thr
 420 425 430
 Asp Cys Thr Ser Glu Gly Arg Arg Asp Asn Met Lys Trp Cys Gly Thr
 435 440 445
 Thr Gln Asn Tyr Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro Met Ala
 450 455 460
 Ala His Glu Glu Ile Cys Thr Thr Asn Glu Gly Val Met Tyr Arg Ile
 465 470 475 480
 Gly Asp Gln Trp Asp Lys Gln His Asp Met Gly His Met Met Arg Cys
 485 490 495
 Thr Cys Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Tyr Ala Tyr Ser
 500 505 510
 Gln Leu Arg Asp Gln Cys Ile Val Asp Asp Ile Thr Tyr Asn Val Asn
 515 520 525
 Asp Thr Phe His Lys Arg His Glu Glu Gly His Met Leu Asn Cys Thr
 530 535 540
 Cys Phe Gly Gln Gly Arg Gly Arg Trp Lys Cys Asp Pro Val Asp Gln
 545 550 555 560
 Cys Gln Asp Ser Glu Thr Gly Thr Phe Tyr Gln Ile Gly Asp Ser Trp
 565 570 575
 Glu Lys Tyr Val His Gly Val Arg Tyr Gln Cys Tyr Cys Tyr Gly Arg
 580 585 590
 Gly Ile Gly Glu Trp His Cys Gln Pro Leu Gln Thr Tyr Pro Ser Ser
 595 600 605
 Ser Gly Pro Val Glu Val Phe Ile Thr Glu Thr Pro Ser Gln Pro Asn
 610 615 620
 Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His Ile Ser Lys
 625 630 635 640
 Tyr Ile Leu Arg Trp Arg Pro Val Ser Ile Pro Pro Arg Asn Leu Gly
 645 650 655
 Tyr

<210> 37
<211> 719
<212> PRT
<213> *Homo sapiens*

<220>
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<222> 676, 679, 683, 717
<223> xaa = Any Amino Acid

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 Leu Asn Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Leu Ala Val
 20 25 30
 Leu Cys Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys
 35 40 45
 Arg Gln Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser
 50 55 60
 Gln Ser Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn
 65 70 75 80
 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys
 85 90 95
 Tyr Gly Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu
 100 105 110
 Glu Thr Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp

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115	120	125
Thr Tyr Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile		
130	135	140
Gly Ala Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His		
145	150	155
Glu Gly Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His		160
165	170	175
Glu Thr Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys		
180	185	190
Gly Glu Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala		
195	200	205
Ala Gly Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln		
210	215	220
Gly Trp Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg		
225	230	235
Ile Thr Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr		240
245	250	255
Ser Tyr Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn		
260	265	270
Leu Leu Gln Cys Ile Cys Thr Gly Asn Gly Arg Gly Glu Trp Lys Cys		
275	280	285
Glu Arg His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe		
290	295	300
Thr Asp Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro		
305	310	315
Pro Pro Tyr Gly His Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val		320
325	330	335
Gly Met Gln Trp Leu Lys Thr Gln Gly Asn Lys Gln Met Leu Cys Thr		
340	345	350
Cys Leu Gly Asn Gly Val Ser Cys Gln Glu Thr Ala Val Thr Gln Thr		
355	360	365
Tyr Gly Gly Asn Ser Asn Gly Glu Pro Cys Val Leu Pro Phe Thr Tyr		
370	375	380
Asn Gly Arg Thr Phe Tyr Ser Cys Thr Thr Glu Gly Arg Gln Asp Gly		
385	390	395
His Leu Trp Cys Ser Thr Thr Ser Asn Tyr Glu Gln Asp Gln Lys Tyr		400
405	410	415
Ser Phe Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn		
420	425	430
Ser Asn Gly Ala Leu Cys His Phe Pro Phe Leu Tyr Asn Asn His Asn		
435	440	445
Tyr Thr Asp Cys Thr Ser Glu Gly Arg Arg Asp Asn Met Lys Trp Cys		
450	455	460
Gly Thr Thr Gln Asn Tyr Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro		
465	470	475
Met Ala Ala His Glu Ile Cys Thr Thr Asn Glu Gly Val Met Tyr		480
485	490	495
Arg Ile Gly Asp Gln Trp Asp Lys Gln His Asp Met Gly His Met Met		
500	505	510
Arg Cys Thr Cys Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Tyr Ala		
515	520	525
Tyr Ser Gln Leu Arg Asp Gln Cys Ile Val Asp Asp Ile Thr Tyr Asn		
530	535	540
Val Asn Asp Thr Phe His Lys Arg His Glu Glu Gly His Met Leu Asn		
545	550	555
Cys Thr Cys Phe Gly Gln Gly Arg Gly Arg Trp Lys Cys Asp Pro Val		
565	570	575
Asp Gln Cys Gln Asp Ser Glu Thr Gly Thr Phe Tyr Gln Ile Gly Asp		
580	585	590
Ser Trp Glu Lys Tyr Val His Gly Val Arg Tyr Gln Cys Tyr Cys Tyr		
595	600	605
Gly Arg Gly Ile Gly Glu Trp His Cys Gln Pro Leu Gln Thr Tyr Pro		
610	615	620

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Ser Ser Ser Gly Pro Val Glu Val Phe Ile Thr Glu Thr Pro Ser Gln
625 630 635 640
Pro Asn Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His Ile
645 650 655
Ser Lys Tyr Ile Leu Arg Trp Arg Pro Val Ser Ile Pro Pro Arg Asn
660 665 670
Leu Gly Tyr Xaa Val Ser Xaa Ser Gln Phe Xaa Trp Phe Leu Phe Phe
675 680 685
Pro Ala Phe Glu Pro Thr Thr Leu Ile Asn Tyr Ser Tyr Ser Ile Tyr
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Tyr Ile Cys Leu Val Asn Lys Gln Tyr Val Val Asn Xaa Ile Asp
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<210> 38
<211> 44
<212> PRT
<213> Homo sapiens

<400> 38
Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp Thr Tyr
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Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile Gly Ala
20 25 30
Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg
35 40

<210> 39
<211> 45
<212> PRT
<213> Homo sapiens

<400> 39
Cys His Glu Gly Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg
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Pro His Glu Thr Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn
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Gly Lys Gly Glu Trp Thr Cys Lys Pro Ile Ala Glu Lys
35 40 45

<210> 40
<211> 45
<212> PRT
<213> Homo sapiens

<400> 40
Cys Phe Asp His Ala Ala Gly Thr Ser Tyr Val Val Gly Glu Thr Trp
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Glu Lys Pro Tyr Gln Gly Trp Met Met Val Asp Cys Thr Cys Leu Gly
20 25 30
Glu Gly Ser Gly Arg Ile Thr Gly Thr Ser Arg Asn Arg
35 40 45

<210> 41
<211> 1926
<212> DNA
<213> Homo sapiens

<400> 41
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atcaactgaga	ctccgagtca	gccccactcc	caccccatcc	agtggaaatgc	accacagcca	1860
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ggatac						1926

<210> 42

<211> 21

<212> PRT

<213> Homo sapiens

<400> 42

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Arg	Asp	Gln	Cys	Ile											
				20											

<210> 43

<211> 20

<212> PRT

<213> Homo sapiens

<400> 43

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Trp	Lys	Glu	Ala												
			20												

<210> 44

<211> 720

<212> PRT

<213> Homo sapiens

<400> 44

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Asn Leu Val Ala Thr Cys Leu Pro Val Arg Ala Ser Leu Pro His Arg
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 Leu Asn Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Ala Val
 20 25 30
 Leu Cys Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys
 35 40 45
 Arg Gln Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser
 50 55 60
 Gln Ser Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn
 65 70 75 80
 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys
 85 90 95
 Tyr Gly Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu
 100 105 110
 Glu Thr Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp
 115 120 125
 Thr Tyr Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile
 130 135 140
 Gly Ala Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His
 145 150 155 160
 Glu Gly Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His
 165 170 175
 Glu Thr Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys
 180 185 190
 Gly Glu Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala
 195 200 205
 Ala Gly Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln
 210 215 220
 Gly Trp Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg
 225 230 235 240
 Ile Thr Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr
 245 250 255
 Ser Tyr Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn
 260 265 270
 Leu Leu Gln Cys Ile Cys Thr Gly Asn Gly Arg Gly Glu Trp Lys Cys
 275 280 285
 Glu Arg His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe
 290 295 300
 Thr Asp Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro
 305 310 315 320
 Pro Pro Tyr Gly His Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val
 325 330 335
 Gly Met Gln Trp Leu Lys Thr Gln Gly Asn Lys Gln Met Leu Cys Thr
 340 345 350
 Cys Leu Gly Asn Gly Val Ser Cys Gln Glu Thr Ala Val Thr Gln Thr
 355 360 365
 Tyr Gly Gly Asn Ser Asn Gly Glu Pro Cys Val Leu Pro Phe Thr Tyr
 370 375 380
 Asn Gly Arg Thr Phe Tyr Ser Cys Thr Thr Glu Gly Arg Gln Asp Gly
 385 390 395 400
 His Leu Trp Cys Ser Thr Thr Ser Asn Tyr Glu Gln Asp Gln Lys Tyr
 405 410 415
 Ser Phe Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn
 420 425 430
 Ser Asn Gly Ala Leu Cys His Phe Pro Phe Leu Tyr Asn Asn His Asn
 435 440 445
 Tyr Thr Asp Cys Thr Ser Glu Gly Arg Arg Asp Asn Met Lys Trp Cys
 450 455 460
 Gly Thr Thr Gln Asn Tyr Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro
 465 470 475 480
 Met Ala Ala His Glu Glu Ile Cys Thr Thr Asn Glu Gly Val Met Tyr
 485 490 495
 Arg Ile Gly Asp Gln Trp Asp Lys Gln His Asp Met Gly His Met Met

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Arg Cys Thr Cys Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Tyr Ala	500	505	510
515	520	525	
Tyr Ser Gln Leu Arg Asp Gln Cys Ile Val Asp Asp Ile Thr Tyr Asn	530	535	540
Val Asn Asp Thr Phe His Lys Arg His Glu Glu Gly His Met Leu Asn	545	550	555
Cys Thr Cys Phe Gly Gln Gly Arg Gly Arg Trp Lys Cys Asp Pro Val	565	570	575
Asp Gln Cys Gln Asp Ser Glu Thr Gly Thr Phe Tyr Gln Ile Gly Asp	580	585	590
Ser Trp Glu Lys Tyr Val His Gly Val Arg Tyr Gln Cys Tyr Cys Tyr	595	600	605
Gly Arg Gly Ile Gly Glu Trp His Cys Gln Pro Leu Gln Thr Tyr Pro	610	615	620
Ser Ser Ser Gly Pro Val Glu Val Phe Ile Thr Glu Thr Pro Ser Gln	625	630	635
640			
Pro Asn Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His Ile	645	650	655
Ser Lys Tyr Ile Leu Arg Trp Arg Pro Lys Asn Ser Val Gly Arg Trp	660	665	670
Lys Glu Ala Thr Ile Pro Gly His Leu Asn Ser Tyr Thr Ile Lys Gly	675	680	685
Leu Lys Pro Gly Val Val Tyr Glu Gly Gln Leu Ile Ser Ile Gln Gln	690	695	700
Tyr Gly His Gln Glu Val Thr Arg Phe Asp Phe Thr Thr Thr Ser Thr	705	710	715
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<210> 45

<211> 11

<212> PRT

<213> Homo sapiens

<400> 45

Thr Ala Ser Gly Val Ala Glu Thr Thr Asn Cys
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